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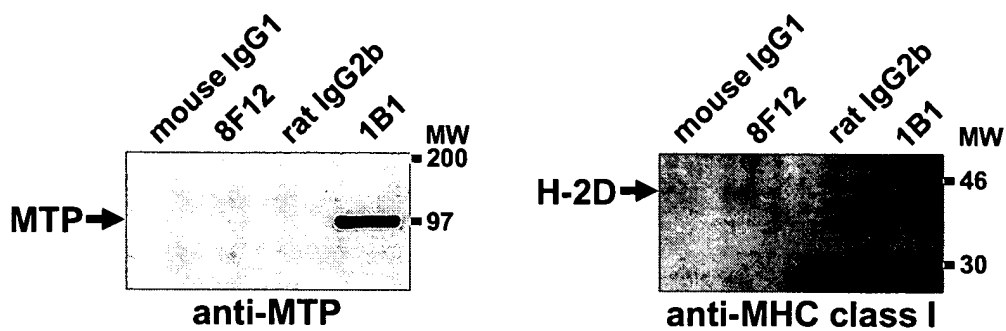
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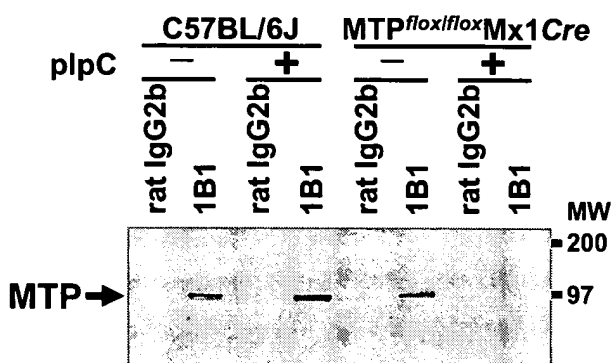
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Figure 1

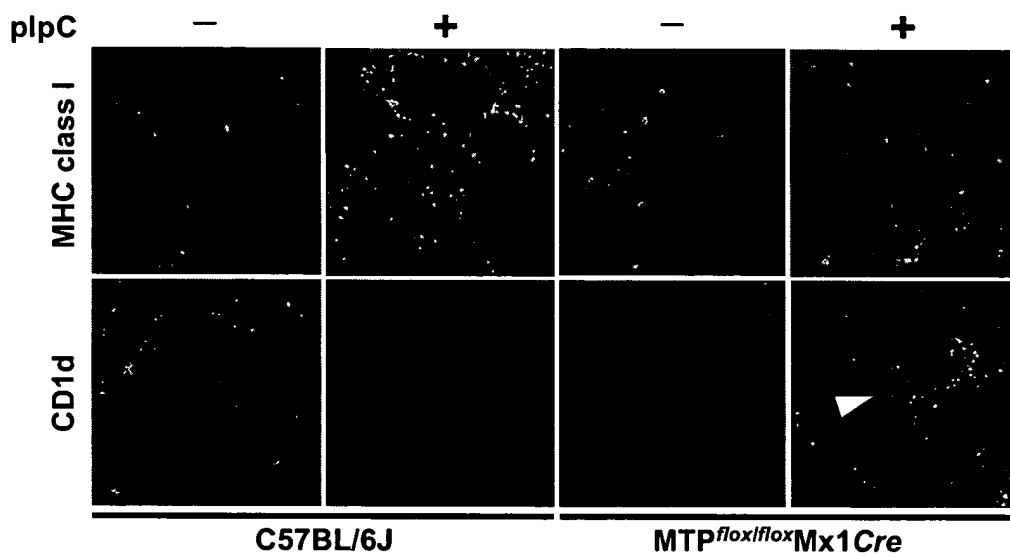
A



B



C



D

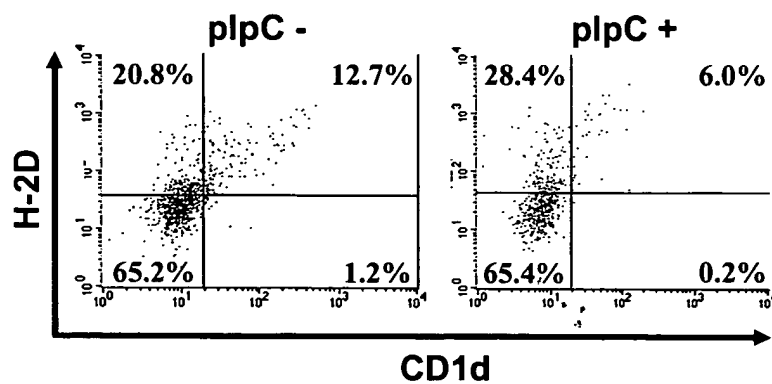


Figure 2

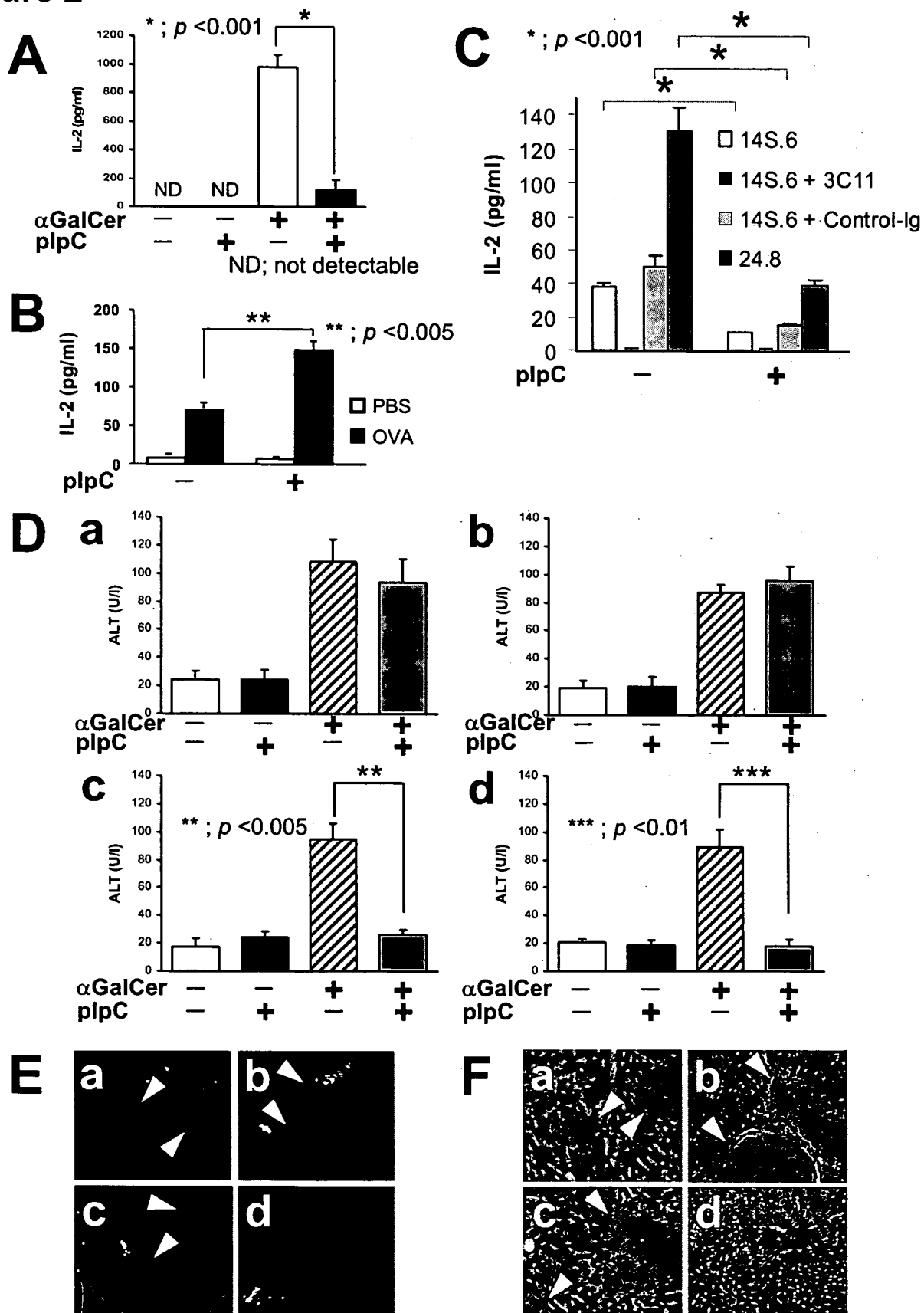
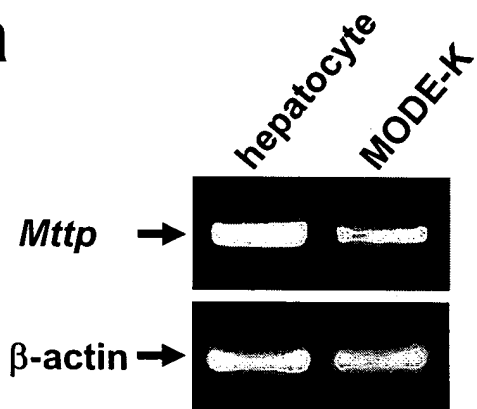
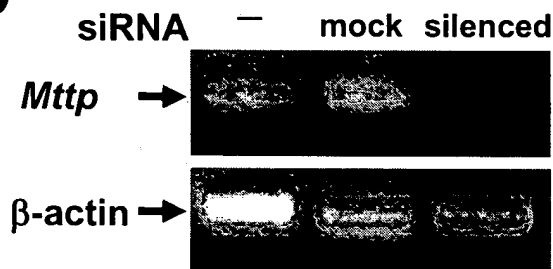


Figure 3

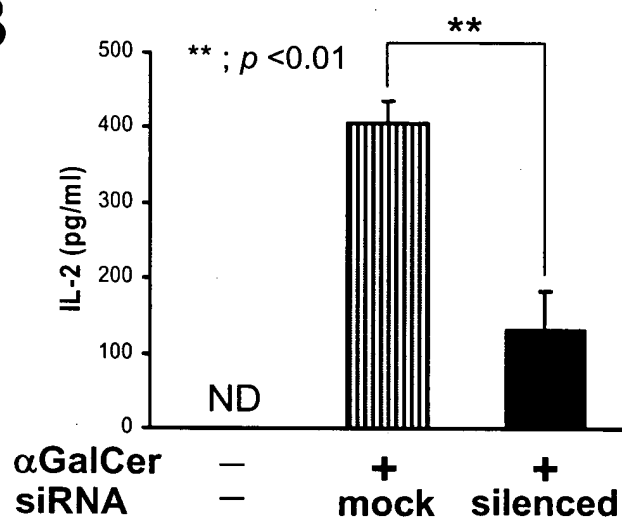
A a



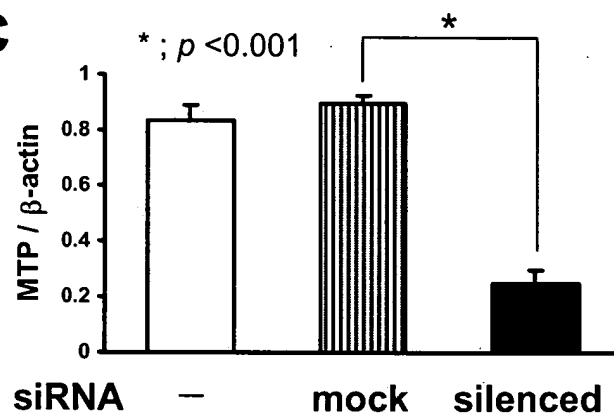
b



B



C



ND; not detectable

Figure 4

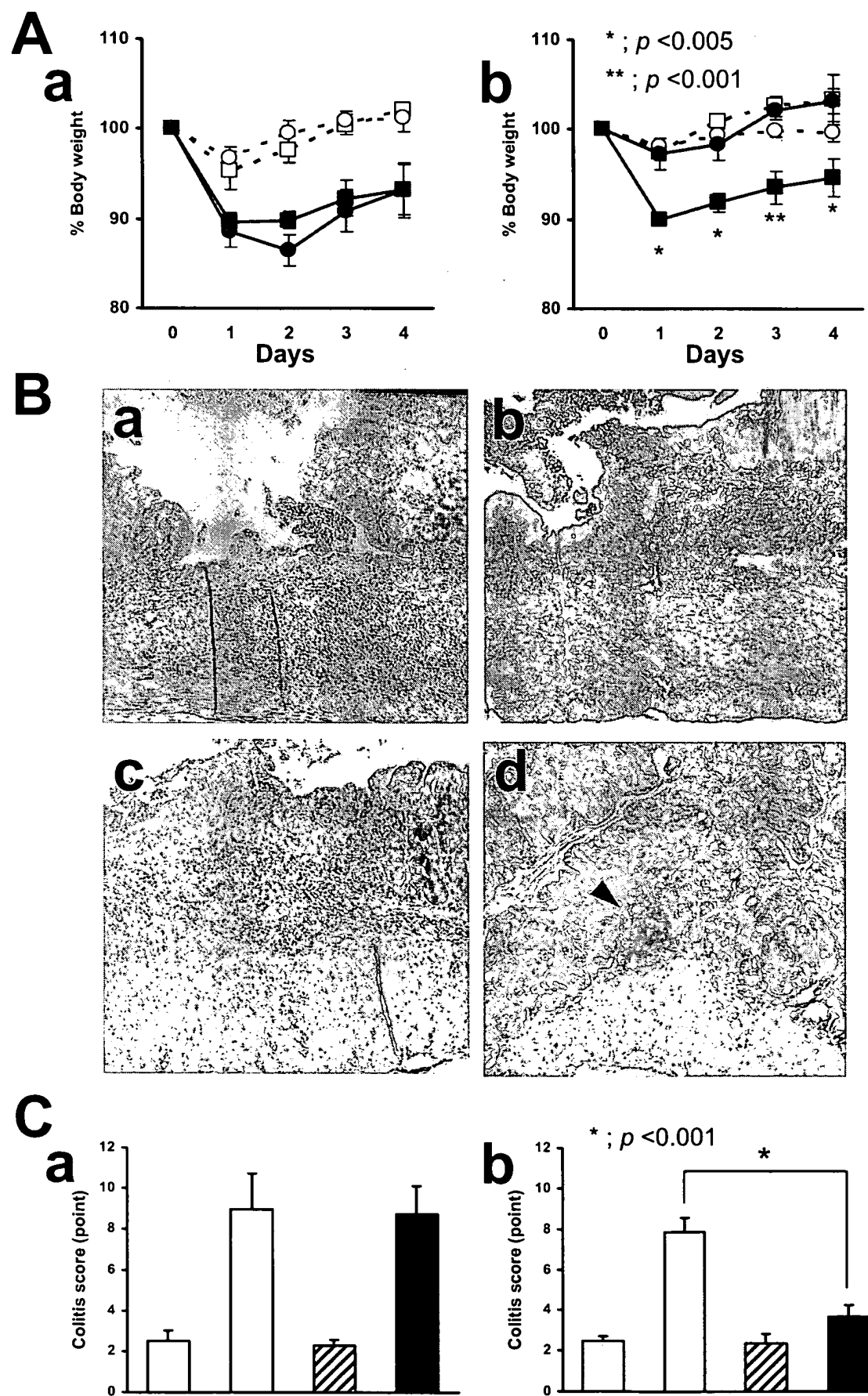
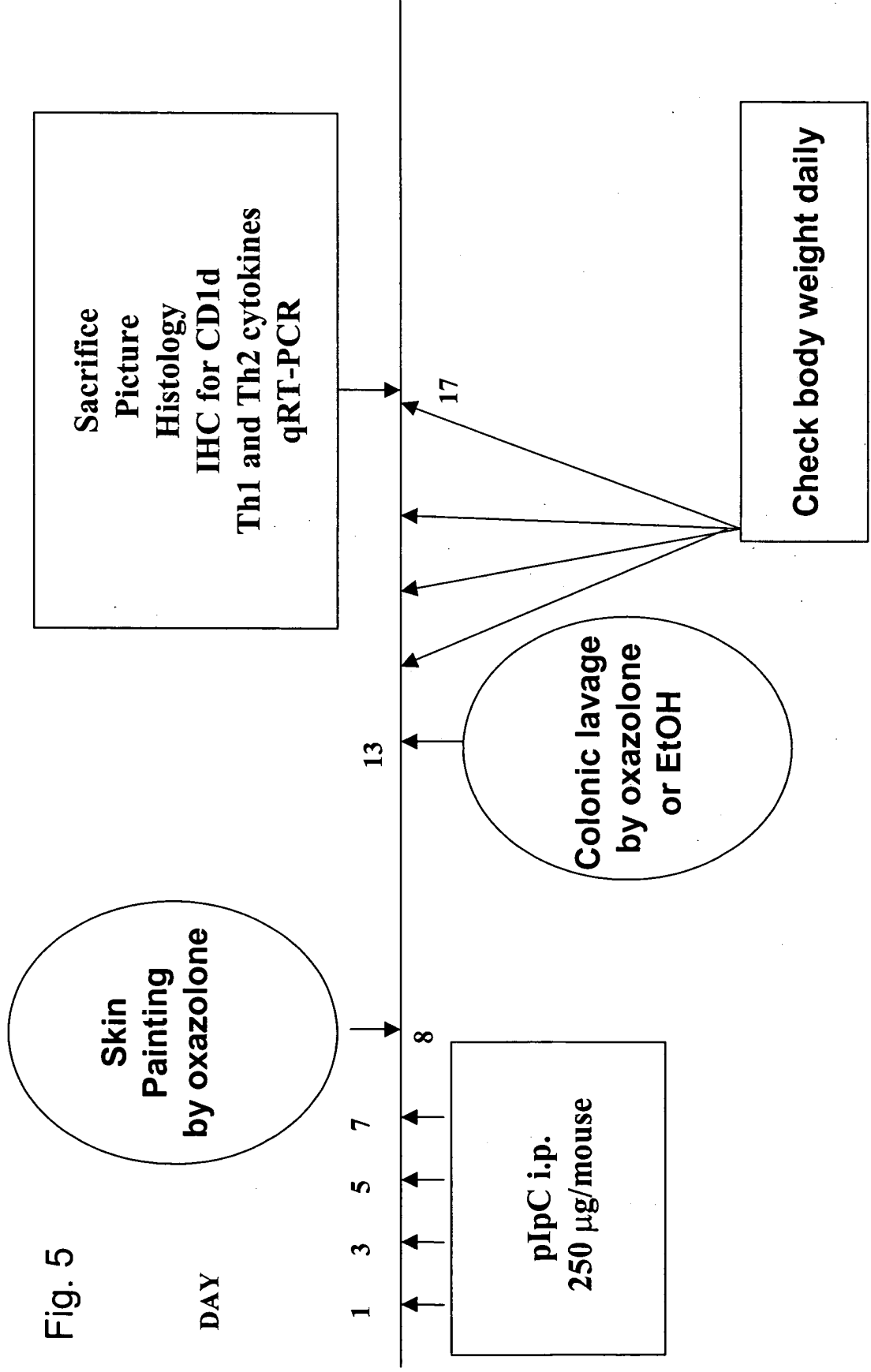


Fig. 5



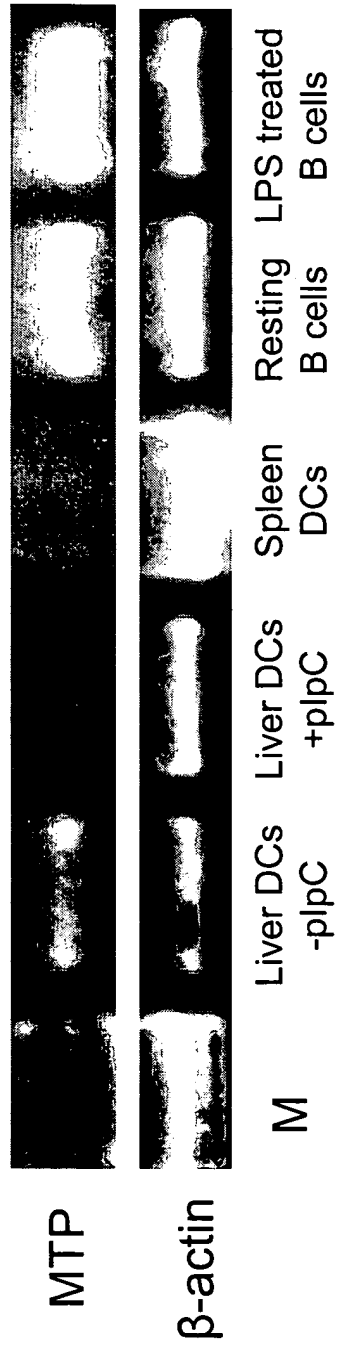


Figure 6

Fig. 7

Silenced and wild type U937 cells are equally capable of presenting alpha gal to NKT cells at a 1:1 effector to target ratio.
U937/DN32.D3 coculture assay
(E:T = 25,000:25,000)



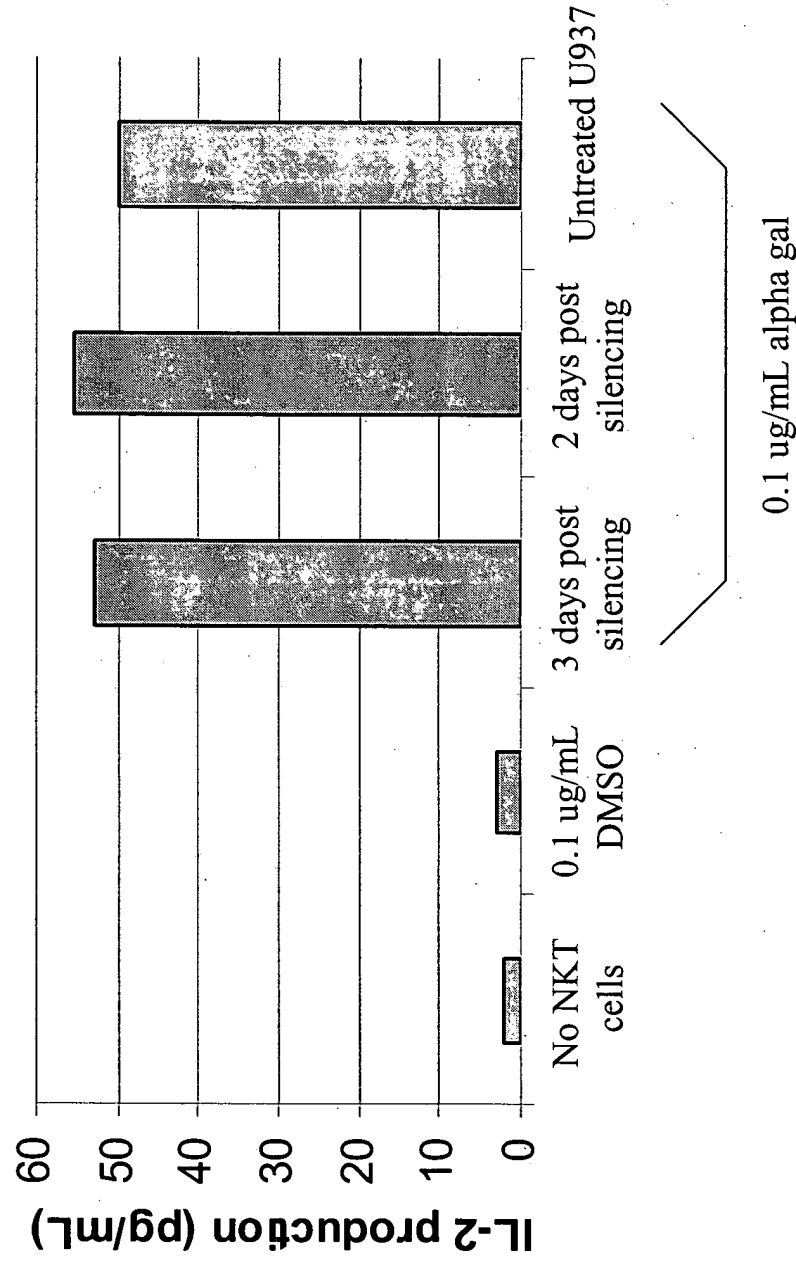
Two days post silencing
 Three days post silencing
 Silenced cells coculture
 Untreated T937

Each data point represents 4 independent trials.
 IL-2 production in the absence of alpha gal was negligible.

Fig.8

Silenced and wild type U937 cells are equally capable of presenting alpha gal to NKT cells at a 10:1 effector to target ratio.

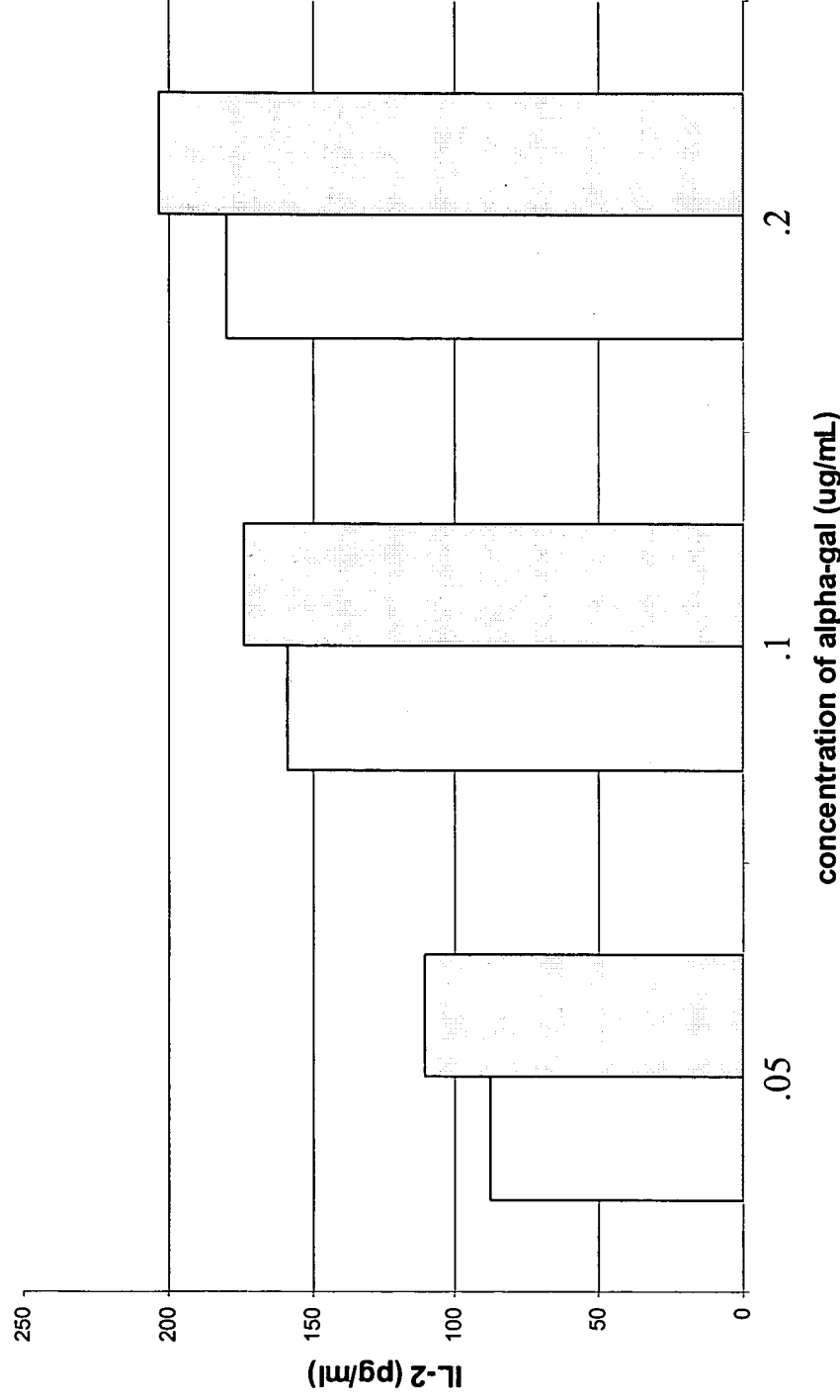
U937/DN32.D3 coculture assay
(E:T = 50,000:5,000)



Each data point represents 4 independent trials.

Fig. 9

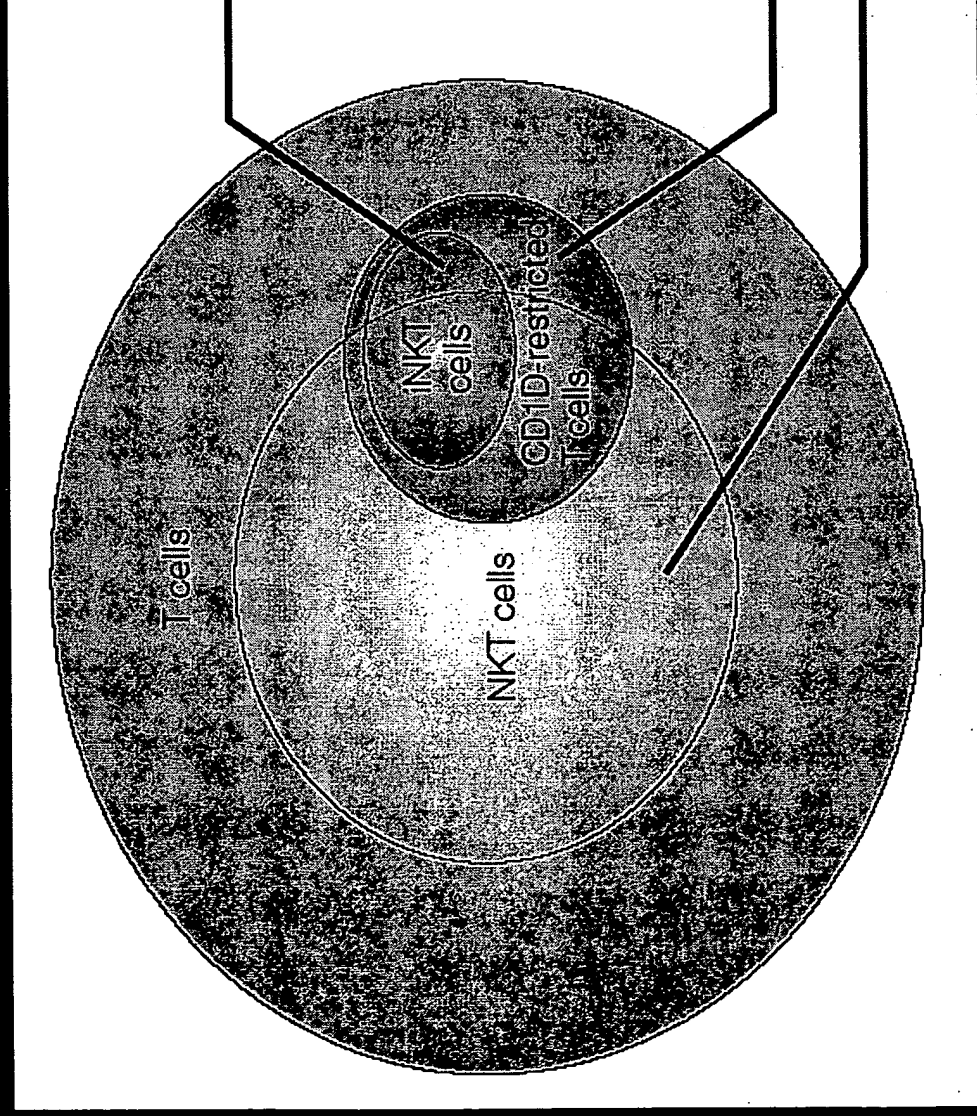
Silenced and wild type U937 cells are not equally capable of presenting alpha gal to NKT cells at a 100:1 effector to target ratio.
U937/DN32.D3 coculture assay
(E:T = 1,000,000:10,000)



Each data point represents one trial.
IL-2 production in the absence of alpha gal was an average of 48.7 pg/mL.

The World of Natural Killer T (NKT) cells:

CD1d-restricted and unrestricted



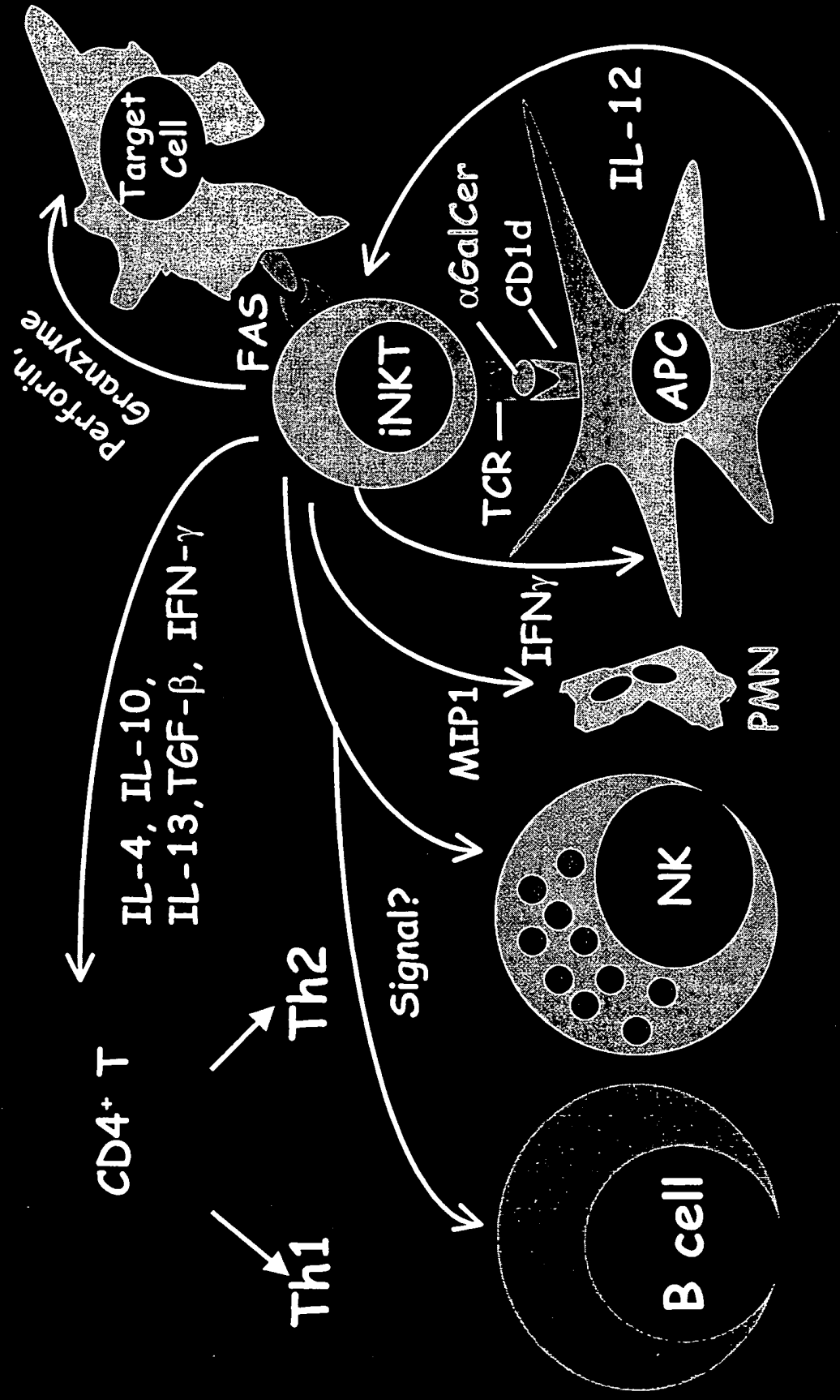
CD4 or DN
V α 14-J α 18
V β 8.2/7/2
 α GalCer

CD4 or DN
Semi-Diverse
V α 3.2-J α 9
V α 8, V β 8

Diverse
CD4, CD8, DN
MHC class I/II

Adapted from Nature Reviews Immunol 2003;3:211

Multiple Regulatory Functions of iNKT Cells



Adapted from Nature Reviews Immunol 2003;3:21

Figure 11

		Section 1				
	(1)	1	10	20	30	40
ApoB 1-269	(1)	-----MDPPRPALLALLA				P LLLLLLAGAR EEE LE
CD1d human	(1)	MGCLLFLLLWALLQAWGSAEVPQRLFP LRCLO				S FANSSWTRTD LAW GE
		Section 2				
	(53)	53	60	70	80	90
ApoB 1-269	(34)	NV	LVC PKD	TRF	HL	K YN-----YEAE--S-SSGVPGT
CD1d human	(53)	LQ	HSWSND	DTV	SL	P QGTFSDQQWETLQHFVRVYRSSFTRDV
		Section 3				
	(105)	105	110	120	130	140
ApoB 1-269	(73)	ATR	NCK	ELEV PQLCSFILK--	S-----	QCTLKEVYGFNP
CD1d human	(105)	MLR	SYP	ELQVSAGCEVHPGNA	NNFFHVA	FQGDILSFQGTSWEPTQEAR
		Section 4				
	(157)	157	170	180	190	208
ApoB 1-269	(108)	EGKALLKKTNS	E	AAAMSR	ELKLAIPEGKQVFLYPEKDEPTYILN	IKRG
CD1d human	(157)	LWVNLAIQVLNQ	K	TRETVO	LLNGTCPQFVSGLLESGKSE----	LKKQV
		Section 5				
	(209)	209	220	230	240	250
ApoB 1-269	(160)	IISALLVPPE	EEAKQ	LF	DTVYGNCSTHFTVKTR	GN-----A-
CD1d human	(204)	KPKAWLSRGP	PGPGR	LL	CHVSGFYKPKPVWVKWM	GEQEQQGTQPGD LP
		Section 6				
	(261)	261	270	280	290	300
ApoB 1-269	(201)	-----T-----		STERDLGQCDRFKPIRTGISP	ALIKGM	RPLSTLI S
CD1d human	(256)	NADETWYLRATL		VAGEAAGLSCRVKHSSLEGQD	VLYWGG	YTSMGLI L
		Section 7				
	(313)	313	320	330	343	
ApoB 1-269	(242)	QS-----CQ		TLDAKRKHVAE		CKEQ
CD1d human	(308)	VLACLLFLLIVG		TSRFKRQTSYQ		L---

Figure 12

		Section 1					
		(1) 1	10	20	30	40	52
ApoB 512-721	(1)	-----					K
CD1d human	(1)	MGCLLFLLWALLQAWGSAEVPQRLFPLRCLQISSFANSSWTRTDGLAWLGE					
		Section 2					
		(53) 53	60	70	80	90	104
ApoB 512-721	(2)	CVQS	KPSLMIQKA	QALRKMEPKDKDQEVLLQTFLDDASPGD	R	AA	LM
CD1d human	(53)	LQTH	WSNDSDTVR	KPWSQGTTFSDQQWETLQHI	FRVYRSSFT	D	KEAK
		Section 3					
		(105) 105	110	120	130	140	158
ApoB 512-721	(54)	RS--	PSQAD	NKIVQ	LPWEQNE	-----QVKN	VAHIAN
CD1d human	(105)	RLSYPLELQ	SAGCE	HPGNASNNFFHVA	FQGKDILSFQGTS	EP	QEAP
		Section 4					
		(157) 157	170	180	190	208	
ApoB 512-721	(89)	LNSEE	D	QDLKKLV	EA	KESQLPTVMDFRKFSRN	-----YQLYKSVL
CD1d human	(157)	WVNLA	Q	LNQDKWT	ET	QWLLNGTCPQFVSGLL	ESGKSELKKQVKFKW
		Section 5					
		(209) 209	220	230	240	250	260
ApoB 512-721	(135)	PSLDP	S	KIEGNL	FDPNN	LPK--ESM	TTLTAFGFASD
CD1d human	(209)	LSRGP	P	PGRLLL	CHVSG	YPKPVWVKW	GEQEQQGTQPD
		Section 6					
		(281) 281	270	280	290	300	312
ApoB 512-721	(184)	KGFEPTL	A	FGKQ	-----FFP	SVNKALYW	-----
CD1d human	(281)	WYLRATL	V	AGEA	GLSCRVKHSSL	GQDIVLYWGGSYTSMGLIALAVLAC	
		Section 7					
		(313) 313	320	335			
ApoB 512-721	(211)	-----					
CD1d human	(313)	LLFLLIVGFTSRFRQTSYQGV					

Figure 13

		Section 1									
		(1)	1	10	20	30	40	52			
ApoB 270-570	(1)	-----FSYNNKYGMVAQVTQTLKLEDTPKINSRFFGEGT						KMGLA	EST		
CD1d human	(1)	MGCLLFLLLWALLQAWGSAEVPQRLFPLRCLQISSFANSSWT						TDGLA	LGE		
		Section 2									
		(53)	53	60	70	80	90	104			
ApoB 270-570	(45)	KSTSPPK	AEAV		LQELKKLTIS	Q	IQRA	LFN	VTELRLGLSDBAVTS		
CD1d human	(53)	LQTHSWS	DSDT		LKPWSQGTFS	Q	WETL	HIF	YRSSFTRDVMEFAK		
		Section 3									
		(105)	105	110	120	130	140	156			
ApoB 270-570	(97)	LPQLIE	SSP		LQALVQCG	PQCSTHI	Q	LK	VHANPLLIDVVTYLVAL		
CD1d human	(105)	LRLSYP	ELQ		AGCEVHPG	ASNFFH	A	QG	DILSFQGT	SWEPTQEAP	
		Section 4									
		(157)	157	170	180	190	208				
ApoB 270-570	(149)	PEPSAQQ	REIFNMAR		QSRATLYALSHAVNNYHK	NP	GTQEL	DIAN			
CD1d human	(157)	WVNLAIQ	LNQDKWTR		TVQWLLNGTCPQFVSGLLE	GK	ELKKQ	KPKA			
		Section 5									
		(209)	209	220	230	240	250	260			
ApoB 270-570	(201)	LMEQIQDD	-----C	GDE	TYTL	LR	IGNMGQTMEQLTPELKS	ILK			
CD1d human	(209)	LSRGPS	PGPGRLLLVCHV		GFYPKPVW	KW	RGEQE	QQTQPGDILPN	DET		
		Section 6									
		(261)	261	270	280	290	300	312			
ApoB 270-570	(244)	CVQSTKPS		IQKAAIQ	LRKMEPKDKDQ		LQTFLLDDASPGDKRLAAYLM				
CD1d human	(261)	WYLRATLD		AGEAAGL	CRVKHSSLEGO		LYWGGSYTSMGLIALAVLAC				
		Section 7									
		(313)	313	320	335						
ApoB 270-570	(296)	L	RSPS	-----							
CD1d human	(313)	L	FLLIVGFTSRFKRQTSYQGVL								

Figure 14